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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/694,519

DATE: 11/08/2000

TIME: 07:33:13

Input Set : A:\ES.txt

Output Set: N:\CRF3\11082000\I694519.raw

ENTERED

3 <110> APPLICANT: Isfort, Robert
 4 Sheldon, Russell
 6 <120> TITLE OF INVENTION: Methods for Identifying Compounds for Regulating Muscle Mass or Function
 7 Using Vasoactive Intestinal Peptide Receptors
 9 <130> FILE REFERENCE: 8311
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/694,519
 C--> 11 <141> CURRENT FILING DATE: 2000-10-23
 11 <160> NUMBER OF SEQ ID NOS: 16
 13 <170> SOFTWARE: PatentIn version 3.0
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 457
 17 <212> TYPE: PRT
 18 <213> ORGANISM: homo sapiens;
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 23 1 5 10 15
 25 Gly Ala Leu Ala Trp Ala Leu Gly Pro Ala Gly Gly Gln Ala Ala Arg
 26 20 25 30
 28 Leu Gln Glu Glu Cys Asp Tyr Val Gln Met Ile Glu Val Gln His Lys
 29 35 40 45
 31 Gln Cys Leu Glu Glu Ala Gln Leu Glu Asn Glu Thr Ile Gly Cys Ser
 32 50 55 60
 34 Lys Met Trp Asp Asn Leu Thr Cys Trp Pro Ala Thr Pro Arg Gly Gln
 35 65 70 75 80
 37 Val Val Val Leu Ala Cys Pro Leu Ile Phe Lys Leu Phe Ser Ser Ile
 38 85 90 95
 40 Gln Gly Arg Asn Val Ser Arg Ser Cys Thr Asp Glu Gly Trp Thr His
 41 100 105 110
 43 Leu Glu Pro Gly Pro Tyr Pro Ile Ala Cys Gly Leu Asp Asp Lys Ala
 44 115 120 125
 46 Ala Ser Leu Asp Glu Gln Gln Thr Met Phe Tyr Gly Ser Val Lys Thr
 47 130 135 140
 49 Gly Tyr Thr Ile Gly Tyr Gly Leu Ser Leu Ala Thr Leu Leu Val Ala
 50 145 150 155 160
 52 Thr Ala Ile Leu Ser Leu Phe Arg Lys Leu His Cys Thr Arg Asn Tyr
 53 165 170 175
 55 Ile His Met His Leu Phe Ile Ser Phe Ile Leu Arg Ala Ala Ala Val
 56 180 185 190
 58 Phe Ile Lys Asp Leu Ala Leu Phe Asp Ser Gly Glu Ser Asp Gln Cys
 59 195 200 205
 61 Ser Glu Gly Ser Val Gly Cys Lys Ala Ala Met Val Phe Phe Gln Tyr
 62 210 215 220
 64 Cys Val Met Ala Asn Phe Phe Trp Leu Leu Val Glu Gly Leu Tyr Leu
 65 225 230 235 240
 67 Tyr Thr Leu Leu Ala Val Ser Phe Phe Ser Glu Arg Lys Tyr Phe Trp
 68 245 250 255
 70 Gly Tyr Ile Leu Ile Gly Trp Gly Val Pro Ser Thr Phe Thr Met Val

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71          260          265          270
73 Trp Thr Ile Ala Arg Ile His Phe Glu Asp Tyr Gly Cys Trp Asp Thr
74          275          280          285
76 Ile Asn Ser Ser Leu Trp Trp Ile Ile Lys Gly Pro Ile Leu Thr Ser
77          290          295          300
79 Ile Leu Val Asn Phe Ile Leu Phe Ile Cys Ile Ile Arg Ile Leu Leu
80 305          310          315          320
82 Gln Lys Leu Arg Pro Asp Ile Arg Lys Ser Asp Ser Ser Pro Tyr
83          325          330          335
85 Ser Arg Leu Ala Arg Ser Thr Leu Leu Leu Ile Pro Leu Phe Gly Val
86          340          345          350
88 His Tyr Ile Met Phe Ala Phe Phe Pro Asp Asn Phe Lys Pro Glu Val
89          355          360          365
91 Lys Met Val Phe Glu Leu Val Val Gly Ser Phe Gln Gly Phe Val Val
92          370          375          380
94 Ala Ile Leu Tyr Cys Phe Leu Asn Gly Glu Val Gln Ala Glu Leu Arg
95 385          390          395          400
97 Arg Lys Trp Arg Arg Trp His Leu Gln Gly Val Leu Gly Trp Asn Pro
98          405          410          415
100 Lys Tyr Arg His Pro Ser Gly Gly Ser Asn Gly Ala Thr Cys Ser Thr
101          420          425          430
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114 <400> SEQUENCE: 2
116 Met Arg Pro Pro Ser Pro Leu Pro Ala Arg Trp Leu Cys Val Leu Ala
117 1          5          10          15
119 Gly Ala Leu Ala Trp Ala Leu Gly Pro Ala Gly Gly Gln Ala Ala Arg
120          20          25          30
122 Leu Gln Glu Glu Cys Asp Tyr Val Gln Met Ile Glu Val Gln His Lys
123          35          40          45
125 Gln Cys Leu Glu Glu Ala Gln Leu Glu Asn Glu Thr Ile Gly Cys Ser
126          50          55          60
128 Lys Met Trp Asp Asn Leu Thr Cys Trp Pro Ala Thr Pro Arg Gly Gln
129 65          70          75          80
131 Val Val Val Leu Ala Cys Pro Leu Ile Phe Lys Leu Phe Ser Ser Ile
132          85          90          95
134 Gln Gly Arg Asn Val Ser Arg Ser Cys Thr Asp Glu Gly Trp Thr His
135          100          105          110
137 Leu Glu Pro Gly Pro Tyr Pro Ile Ala Cys Gly Leu Asp Asp Lys Ala
138          115          120          125
140 Ala Ser Leu Asp Glu Gln Gln Thr Met Phe Tyr Gly Ser Val Lys Thr
141          130          135          140
143 Gly Tyr Thr Ile Gly Tyr Gly Leu Ser Leu Ala Thr Leu Leu Val Ala

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144 145          150          155          160
146 Thr Ala Ile Leu Ser Leu Phe Arg Lys Leu His Cys Thr Arg Asn Tyr
147          165          170          175
149 Ile His Met His Leu Phe Ile Ser Phe Ile Leu Arg Ala Ala Val
150          180          185          190
152 Phe Ile Lys Asp Leu Ala Leu Phe Asp Ser Gly Glu Ser Asp Gln Cys
153          195          200          205
155 Ser Glu Gly Ser Val Gly Cys Lys Ala Ala Met Val Phe Phe Gln Tyr
156          210          215          220
158 Cys Val Met Ala Asn Phe Phe Trp Leu Leu Val Glu Gly Leu Tyr Leu
159 225          230          235          240
161 Tyr Thr Leu Leu Ala Val Ser Phe Phe Ser Glu Arg Lys Tyr Phe Trp
162          245          250          255
164 Gly Tyr Ile Leu Ile Gly Trp Gly Val Pro Ser Thr Phe Thr Met Val
165          260          265          270
167 Trp Thr Ile Ala Arg Ile His Phe Glu Asp Tyr Gly Leu Leu Arg Cys
168          275          280          285
170 Trp Asp Thr Ile Asn Ser Ser Leu Trp Trp Ile Ile Lys Gly Pro Ile
171          290          295          300
173 Leu Thr Ser Ile Leu Val Asn Phe Ile Leu Phe Ile Cys Ile Ile Arg
174 305          310          315          320
176 Ile Leu Leu Gln Lys Leu Arg Pro Pro Asp Ile Arg Lys Ser Asp Ser
177          325          330          335
179 Ser Pro Tyr Ser Arg Leu Ala Arg Ser Thr Leu Leu Leu Ile Pro Leu
180          340          345          350
182 Phe Gly Val His Tyr Ile Met Phe Ala Phe Phe Pro Asp Asn Phe Lys
183          355          360          365
185 Pro Glu Val Lys Met Val Phe Glu Leu Val Val Gly Ser Phe Gln Gly
186          370          375          380
188 Phe Val Val Ala Ile Leu Tyr Cys Phe Leu Asn Gly Glu Val Gln Ala
189 385          390          395          400
191 Glu Leu Arg Arg Lys Trp Arg Arg Trp His Leu Gln Gly Val Leu Gly
192          405          410          415
194 Trp Asn Pro Lys Tyr Arg His Pro Ser Gly Gly Ser Asn Gly Ala Thr
195          420          425          430
197 Cys Ser Thr Gln Val Ser Met Leu Thr Arg Val Ser Pro Gly Ala Arg
198          435          440          445
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201          450          455          460
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204 <211> LENGTH: 459
205 <212> TYPE: PRT
206 <213> ORGANISM: rattus norvegicus;
208 <400> SEQUENCE: 3
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211 1          5          10          15
213 Gly Ala Leu Ala Cys Ala Leu Arg Pro Ala Gly Ser Gln Ala Ala Ser
214          20          25          30
216 Pro Gln His Glu Cys Glu Tyr Leu Gln Leu Ile Glu Ile Gln Arg Gln

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217		35		40		45	
219	Gln	Cys	Leu	Glu	Glu	Ala	Gln
220		50		55		60	
222	Lys	Met	Trp	Asp	Asn	Leu	Thr
223	65			70		75	
225	Ala	Val	Val	Leu	Asp	Cys	Pro
226				85		90	
228	His	Gly	Tyr	Asn	Ile	Ser	Arg
229				100		105	
231	Leu	Glu	Pro	Gly	Pro	Tyr	His
232				115		120	
234	Ser	Ser	Leu	Asp	Glu	Gln	Gln
235				130		135	
237	Thr	Gly	Tyr	Thr	Ile	Gly	Tyr
238	145			150		155	
240	Ala	Met	Ala	Ile	Leu	Ser	Leu
241				165		170	
243	Tyr	Ile	His	Met	His	Leu	Phe
244				180		185	
246	Val	Phe	Ile	Lys	Asp	Met	Ala
247				195		200	
249	Cys	Ser	Glu	Ala	Ser	Val	Gly
250				210		215	
252	Tyr	Cys	Val	Met	Ala	Asn	Phe
253	225			230		235	
255	Leu	Tyr	Thr	Leu	Leu	Ala	Val
256				245		250	
258	Trp	Gly	Tyr	Ile	Leu	Ile	Gly
259				260		265	
261	Ile	Trp	Thr	Val	Val	Arg	Ile
262				275		280	
264	Thr	Ile	Ile	Asn	Ser	Ser	Leu
265				290		295	
267	Leu	Ser	Ile	Leu	Val	Asn	Phe
268	305			310		315	
270	Leu	Val	Gln	Lys	Leu	Arg	Pro
271				325		330	
273	Pro	Tyr	Ser	Arg	Leu	Ala	Lys
274				340		345	
276	Gly	Ile	His	Tyr	Val	Met	Phe
277				355		360	
279	Gln	Val	Lys	Met	Val	Phe	Glu
280				370		375	
282	Val	Val	Ala	Ile	Leu	Tyr	Cys
283	385			390		395	
285	Leu	Arg	Arg	Lys	Trp	Arg	Arg
286				405		410	
288	Ser	Ser	Lys	Ser	Gln	His	Pro
289				420		425	

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291 Ser Thr Gln Val Ser Met Leu Thr Arg Val Ser Pro Ser Ala Arg Arg
292          435          440          445
294 Ser Ser Ser Phe Gln Ala Glu Val Ser Leu Val.
295    450          455
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298 <211> LENGTH: 459
299 <212> TYPE: PRT
300 <213> ORGANISM: mus musculus;
302 <400> SEQUENCE: 4
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305 1          5          10          15
307 Gly Ala Leu Ala Cys Ala Leu Gly Pro Ala Gly Ser Arg Ala Ala Ser
308          20          25          30
310 Pro His Gln Glu Cys Glu Tyr Leu Gln Met Ile Glu Lys Gln Arg Gln
311          35          40          45
313 Gln Cys Leu Glu Glu Ala Gln Leu Glu Asn Lys Thr Thr Gly Cys Ser
314    50          55          60
316 Lys Met Trp Asp Asn Leu Thr Cys Trp Pro Thr Thr Pro Trp Gly Gln
317 65          70          75          80
319 Val Val Val Leu Asp Cys Pro Leu Ile Phe Gln Leu Phe Ser Pro Ile
320          85          90          95
322 His Gly Tyr Asn Ile Ser Arg Asn Cys Thr Glu Glu Gly Trp Ser Gln
323          100         105         110
325 Leu Glu Pro Gly Pro Tyr His Ile Ala Cys Gly Leu Asn Asp Arg Ala
326    115         120         125
328 Ser Ser Met Asp Glu Gln Gln Thr Glu Phe Tyr Asp Ala Val Lys
329    130         135         140
331 Thr Gly Tyr Thr Ile Gly Tyr Ser Leu Ser Leu Ala Ser Leu Leu Val
332 145         150         155         160
334 Ala Met Ala Ile Leu Ser Leu Phe Arg Lys Leu His Cys Thr Arg Asn
335          165         170         175
337 Tyr Ile His Met His Leu Phe Met Ser Phe Ile Leu Arg Ala Thr Ala
338          180         185         190
340 Val Phe Ile Lys Asp Met Ala Leu Phe Asn Asn Gly Glu Thr Asp His
341          195         200         205
343 Cys Ser Glu Ala Ser Val Ser Cys Lys Ala Ala Val Val Phe Phe Gln
344    210         215         220
346 Tyr Cys Val Met Ala Asn Phe Phe Trp Leu Leu Val Glu Gly Leu Tyr
347 225         230         235         240
349 Leu His Thr Leu Leu Ala Val Ser Phe Phe Ser Glu Arg Lys Tyr Phe
350          245         250         255
352 Trp Gly Tyr Ile Leu Ile Gly Trp Gly Val Pro Ser Val Phe Ile Met
353          260         265         270
355 Ile Trp Thr Ile Val Arg Ile His Phe Glu Asp Phe Gly Cys Trp Asp
356          275         280         285
358 Thr Ile Ile Asn Ser Ser Leu Trp Trp Ile Ile Lys Gly Pro Ile Leu
359    290         295         300
361 Ile Ser Ile Leu Val Asn Phe Ile Leu Phe Ile Cys Ile Ile Arg Ile
362 305          310          315          320

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VERIFICATION SUMMARY DATE: 11/08/2000
PATENT APPLICATION: US/09/694,519 TIME: 07:33:14

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Output Set: N:\CRF3\11082000\I694519.raw

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date